





RA McMurray A.;  
RT "2.2 Mb of contiguous nucleotide sequence from chromosome III of C.  
RT elegans.";  
RL Submitted (JUN-1995) to the EMBL/GenBank/DBJ databases.  
RN [2]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=94150718; PubMed=7906398;  
RA Wilson R., Almscough R., Anderson K., Baynes C., Berks M.,  
RA Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A.,  
RA Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.,  
RA Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.,  
RA Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,  
RA Lightning J., Lloyd C., McMurray A., Mortimore B., O'Callaghan M.,  
RA Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen R.,  
RA Thierly-Mieg J., Thomas K., Vaughan M., Vaughan K., Waterston R.,  
RA Watson A., Weinstock L., Wilkinson-Sprat J., Wohlman P.;  
RT "2.2 Mb of contiguous nucleotide sequence from chromosome III of C.  
RT elegans.";  
RL Nature 368:32-38(1994).  
EMBL: Z49886; CAA90055.1; -.  
DR INTERPRO: IPR000561; -.  
DR INTERPRO: IPR001007; -.  
DR INTERPRO: IPR001450; -.  
DR INTERPRO: IPR001502; -.  
DR PROSITE: PS00022; EGF\_1; UNKNOWN\_1.  
DR PROSITE: PS00198; 4FE4S\_FERRDOXIN; UNKNOWN\_1.  
DR PROSITE: PS00764; ENDONUCLEASE III\_1; 1.  
DR PROSITE: PS01208; VMFC; UNKNOWN\_1.  
SQ SEQUENCE 152 AA; 15645 MW; 6E254F0BE476D354 CRC64;

Query Match 100.0%; Score 75; DB 5; Length 152;  
Best Local Similarity 22.2%; Pred. No. 16;  
Matches 6; Conservative 21; Mismatches 0; Indels 0; Gaps 0;

OY 1 CXCXXXXXXXCCCCXXXXC 27  
Db 88 CTCCTCCCTCCCTCCCTCCCTCCGCGC 114

RESULT 5  
ID Q22048 PRELIMINARY; PRT; 164 AA.  
AC Q22048;  
DT 01-NOV-1996 (TReMBLrel. 01, Created)  
DT 01-NOV-1996 (TReMBLrel. 01, Last sequence update)  
DT 01-MAY-2000 (TReMBLrel. 13, Last annotation update)  
DE T01B7.8 PROTEIN.  
GN T01B7.8.  
OS Caenorhabditis elegans.  
OC Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditioidea;  
OC Rhabditidae; Pelodierinae; Caenorhabditis.  
OX NCBI\_TaxID=6239.  
RN [1]  
RP SEQUENCE FROM N.A.  
RL Sims M.;  
RT Submitted (OCT-1995) to the EMBL/GenBank/DBJ databases.  
RN [2]  
RP SEQUENCE FROM N.A.  
RX MEDLINE=94150718; PubMed=7906398;  
RA Wilson R., Almscough R., Anderson K., Baynes C., Berks M.,  
RA Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A.,  
RA Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.,  
RA Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.,  
RA Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,  
RA Lightning J., Lloyd C., McMurray A., Mortimore B., O'Callaghan M.,  
RA Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen R.,  
RA Shalton N., Smith A., Sonhammer E., Staden R., Sulston J.,  
RA Thierly-Mieg J., Thomas K., Vaughan M., Vaughan K., Waterston R.,  
RA Watson A., Weinstock L., Wilkinson-Sprat J., Wohlman P.;  
RT "2.2 Mb of contiguous nucleotide sequence from chromosome III of C.  
RT elegans.";

RL Nature 368:32-38(1994).  
EMBL: Z66499; CAA91301.1; -.  
DR INTERPRO: IPR000561; -.  
DR INTERPRO: IPR001007; -.  
DR INTERPRO: IPR001271; -.  
DR INTERPRO: IPR001450; -.  
DR INTERPRO: IPR001502; -.  
DR PROSITE: PS00022; EGF\_1; UNKNOWN\_1.  
DR PROSITE: PS00198; 4FE4S\_FERRDOXIN; UNKNOWN\_1.  
DR PROSITE: PS00269; DEFENSIN; UNKNOWN\_1.  
DR PROSITE: PS01208; VMFC; UNKNOWN\_1.  
SQ SEQUENCE 164 AA; 16499 MW; C002D4B36C9FCED CRC64;

Query Match 100.0%; Score 75; DB 5; Length 164;  
Best Local Similarity 22.2%; Pred. No. 17;  
Matches 6; Conservative 21; Mismatches 0; Indels 0; Gaps 0;

OY 1 CXCXXXXXXXCCCCXXXXC 27  
Db 84 CCCCTCCCTCCCTCCCTCCCTCC 110

RESULT 6  
ID Q18238 PRELIMINARY; PRT; 188 AA.  
AC Q18238;  
DT 01-NOV-1996 (TReMBLrel. 01, Created)  
DT 01-NOV-1996 (TReMBLrel. 01, Last sequence update)  
DT 01-OCT-2000 (TReMBLrel. 15, Last annotation update)  
DE COSMID C27A2.  
GN C27A2.5.  
OS Caenorhabditis elegans.  
OC Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditioidea;  
OC Rhabditidae; Pelodierinae; Caenorhabditis.  
OX NCBI\_TaxID=6239.  
RN [1]  
RP SEQUENCE FROM N.A.  
RC STRAIN=BRISTOL N2;  
RX MEDLINE=94150718; PubMed=7906398;  
RA Wilson R., Almscough R., Anderson K., Baynes C., Berks M.,  
RA Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A.,  
RA Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.,  
RA Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.,  
RA Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,  
RA Lightning J., Lloyd C., McMurray A., Mortimore B., O'Callaghan M.,  
RA Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen R.,  
RA Shalton N., Smith A., Sonhammer E., Staden R., Sulston J.,  
RA Thierly-Mieg J., Thomas K., Vaughan M., Vaughan K., Waterston R.,  
RA Watson A., Weinstock L., Wilkinson-Sprat J., Wohlman P.;  
RT "2.2 Mb of contiguous nucleotide sequence from chromosome III of C.  
RT elegans.";  
RL Nature 368:32-38(1994).  
RN [2]  
RP SEQUENCE FROM N.A.  
RC STRAIN=BRISTOL N2;  
RL Nhan M.;  
RT Submitted (MAY-1996) to the EMBL/GenBank/DBJ databases.  
RN [3]  
RP SEQUENCE FROM N.A.  
RC STRAIN=BRISTOL N2;  
RL Waterston R.;  
EMBL: U58760; AAB00710.1; -.  
DR INTERPRO: IPR001007; -.  
DR INTERPRO: IPR001271; -.  
DR INTERPRO: IPR001450; -.  
DR PROSITE: PS00198; 4FE4S\_FERRDOXIN; UNKNOWN\_1.  
DR PROSITE: PS00269; DEFENSIN; 1.  
DR PROSITE: PS01208; VMFC; UNKNOWN\_1.  
SQ SEQUENCE 188 AA; 18878 MW; 0C5D0DC5CA8E0C4B CRC64;

Query Match 100.0%; Score 75; DB 5; Length 188;









